

**REMARKS**

The replacement Formal Drawings for Fig. 8, Figs. 9a and 9b and Fig. 11 correct errors introduced into the drawings when converted from the informal drawings filed with the parent Application No. 08/341,555, filed June 24, 1996 and insert a missing amino acid residue one-letter symbol at the appropriate position. Copies of the informal drawings for Fig. 8, Figs. 9a and 9b and Fig. 11 are included for the convenience of the Examiner.

The corrections to Fig. 8 include a change in numbering for the initial amino acid residue from position 1 to an indication that this residue is an  $\alpha$ -amino-n-butyric acid (Aba) residue. That this residue is Aba is described on page 47, lines 8-10, where the "monomer of the HIV-1 protease (SF2 isolate)" "99 amino acid residue polypeptide chain" is illustrated. Line 10 indicates that "Aba =  $\alpha$ -amino-n-butyric acid". It is also apparent by inspection that this residue occupies position 67 in this polypeptide, rather than position 1 as labeled in the uncorrected Fig. 8. This is supported by the Brief Description of Fig. 8 at page 8, lines 7-9, which indicates that the peptide shown is "in the region 87-67". The "H" residue two residues downstream from this "Aba" residue at position 67 has also been corrected to "69" from "59" in concordance with the sequence on page 47. The unlabeled peak in Fig. 8 has also been corrected to insert the appropriate "G" into the peptide sequence in agreement with the corresponding "G" at position 73 of the sequence on page 47. The corresponding entry in the Sequence Listing (SEQ ID NO:1) includes the omitted "G", or "Gly" residue at this position.

The amended Fig. 9a and Fig. 9b correct the residue numbering of "I" at position 66 and the "G" at position 48, respectively, in agreement with their respective positions in the sequence on page 47.

The amendment to Fig. 11 corrects an error of a typographical nature. The amino terminus should be indicated by "-NH<sub>2</sub>".

Applicants request entry of the amendments to the specification in adherence with 37 C.F.R. §§1.821 to 1.825. The information contained in the computer readable form of the Sequence Listing for Application No. 08/341,555, submitted December 1, 1998, was prepared through the use of the software program "PatentIn" and is identical to that of the paper copy, a

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copy of which is enclosed for the convenience of the Examiner. This amendment contains no new matter.

If the Examiner believes a telephone conference would expedite prosecution of this application, please telephone the undersigned at 415-576-0200.

Respectfully submitted,



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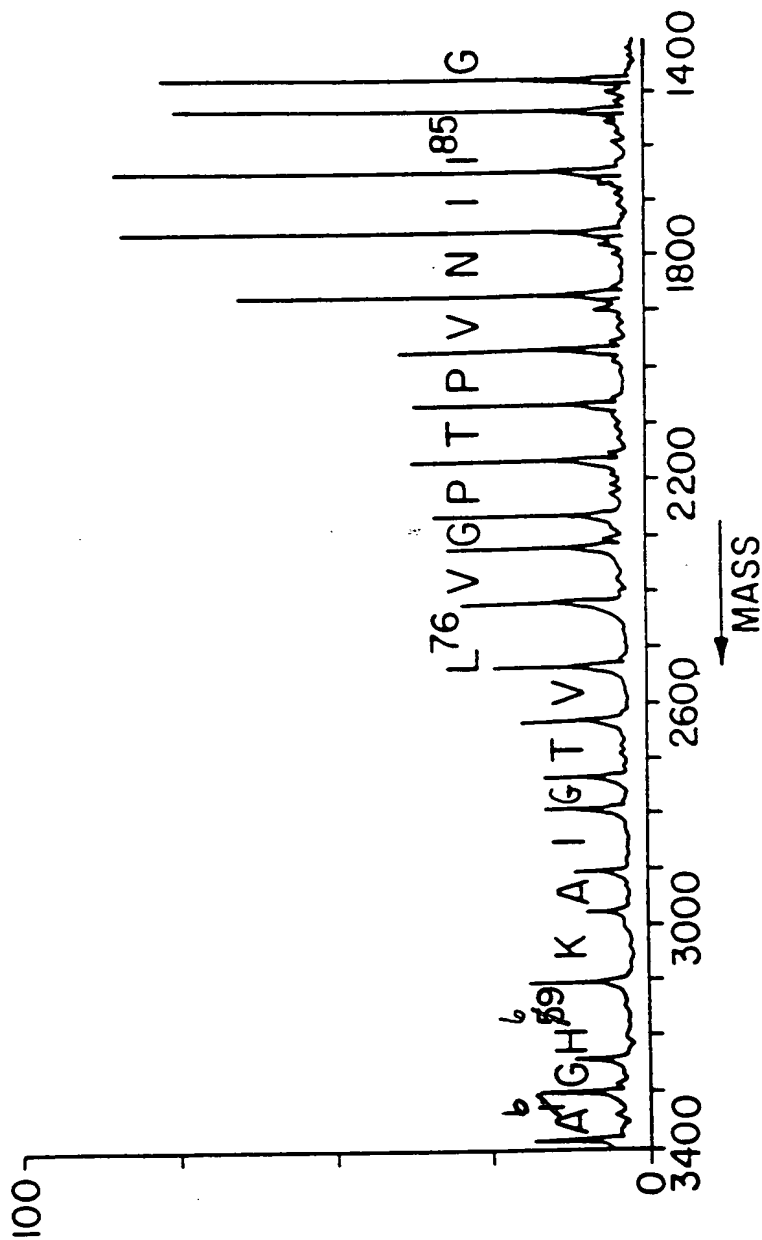


FIG. 8



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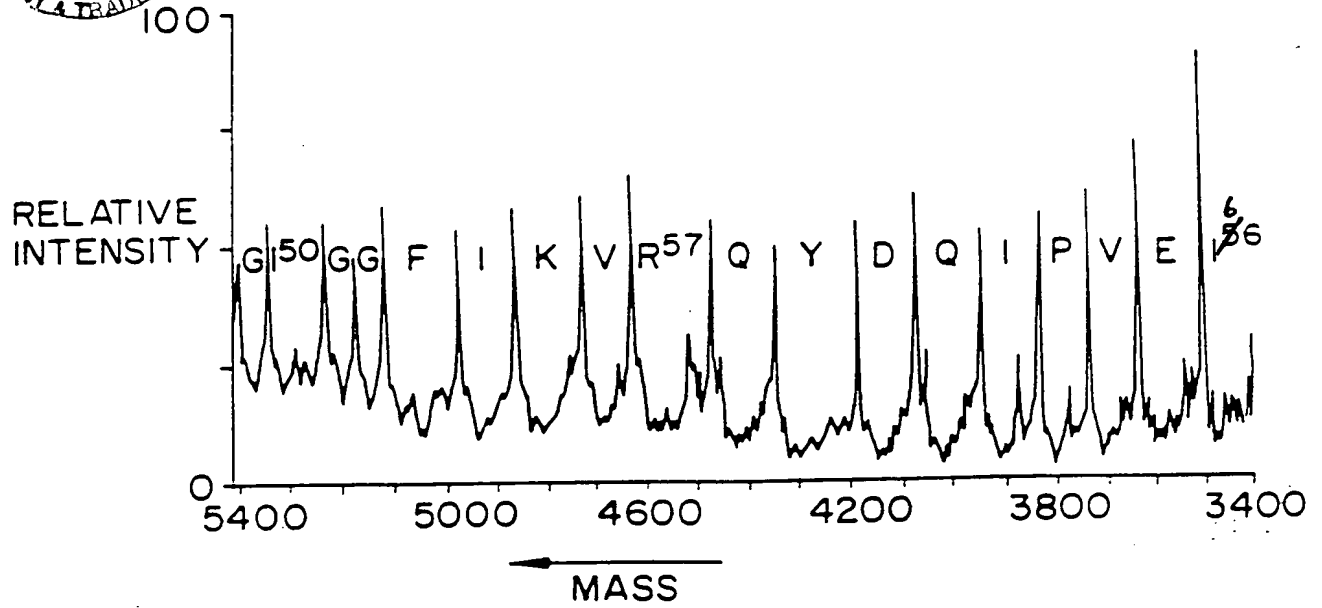


FIG. 9a

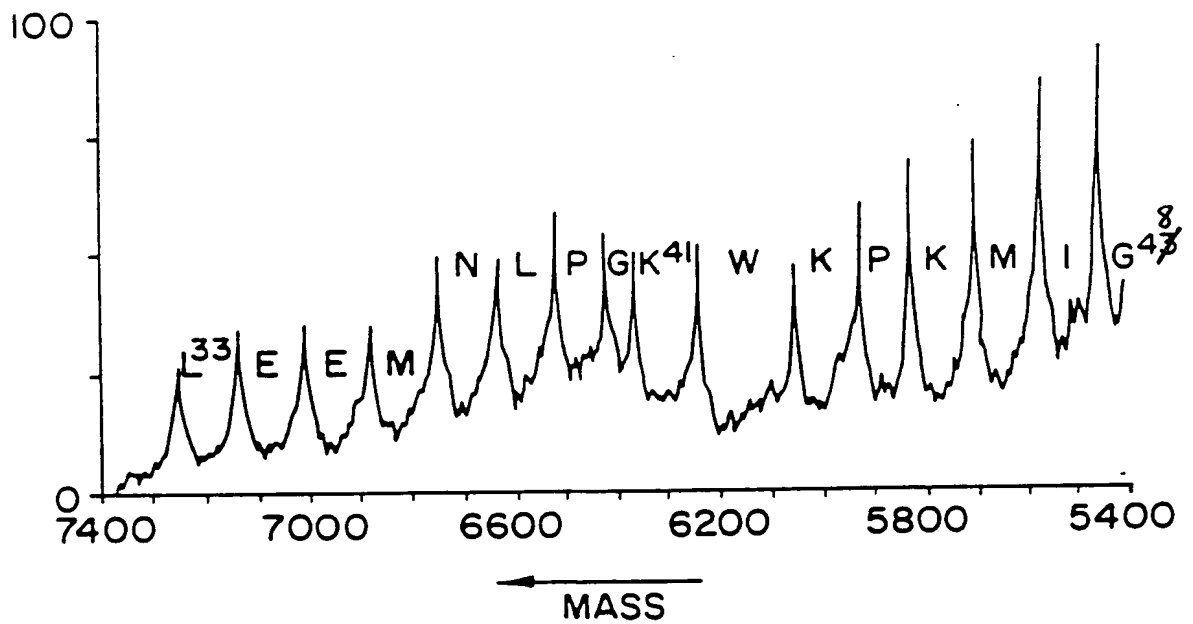


FIG. 9b

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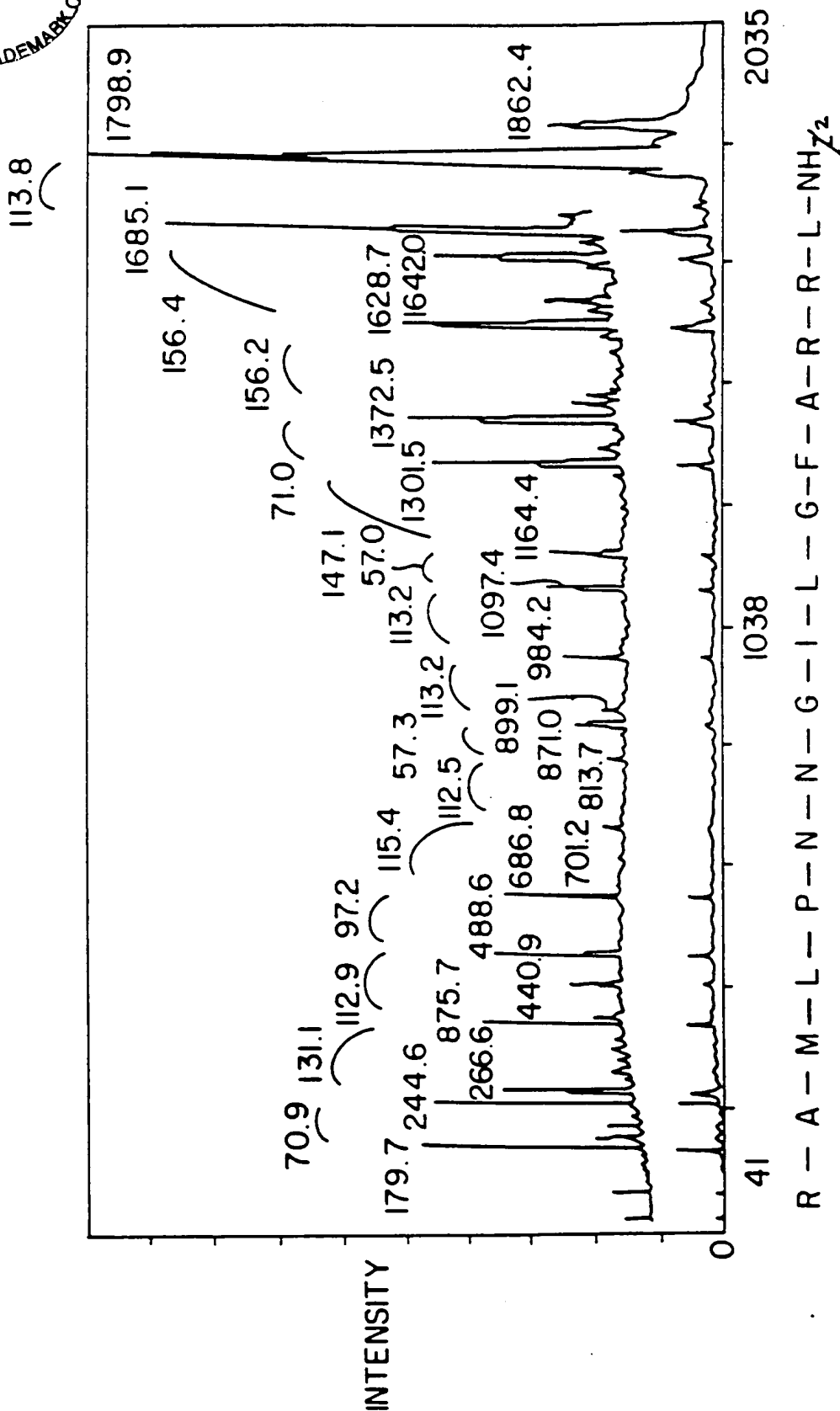


FIG. II